

SEQUENCE LISTING

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<120> INHIBITORS OF ANGIOTENSIN CONVERTING ENZYME

<130> 239568US0CONT

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<151> 2001-10-04

<160> 14

<170> PatentIn version 3.1

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Ser Ser Ala Leu Asp Met Ser Ile Ile Ser Tyr Asp Ser Ala His
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gcg gac aag gcc gcc acg ttg cgc acc gag gag gag ctg atg tcc atg 138
Ala Asp Lys Ala Ala Thr Leu Arg Thr Glu Glu Glu Leu Met Ser Met
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tac gag cag tgg ctc gtg aag cac ggg aag gtg tac aac gcg ctc ggc 186
Tyr Glu Gln Trp Leu Val Lys His Gly Lys Val Tyr Asn Ala Leu Gly

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-75

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| gag aag gag aag cgc ttc cag atc ttc aag gac aac ctg cga ttc atc Glu Lys Glu Lys Arg Phe Gln Ile Phe Lys Asp Asn Leu Arg Phe Ile -70 -65 -60 -55 | 234 |
| gac gac cac aac tcc gcg gag gac cga acc tac aag ctc gga ctg aac Asp Asp His Asn Ser Ala Glu Asp Arg Thr Tyr Lys Leu Gly Leu Asn -50 -45 -40 | 282 |
| cgg ttc gct gat ctc acc aac gag gaa tac agg gcc aag tac ttg gga Arg Phe Ala Asp Leu Thr Asn Glu Glu Tyr Arg Ala Lys Tyr Leu Gly -35 -30 -25 | 330 |
| acc aag atc gat ccc aac cgg agg ctc gga aag acc ccg agc aac cgc Thr Lys Ile Asp Pro Asn Arg Arg Leu Gly Lys Thr Pro Ser Asn Arg -20 -15 -10 | 378 |
| tac gcg cca cgt gtc ggc gac aaa ttg cct gat tcc gtt gat tgg agg Tyr Ala Pro Arg Val Gly Asp Lys Leu Pro Asp Ser Val Asp Trp Arg -5 -1 1 5 10 | 426 |
| aag gaa ggt gct gtt cct ctc gtc aaa gac caa gga ggc tgg ggg agc Lys Glu Gly Ala Val Pro Pro Val Lys Asp Gln Gly Gly Cys Gly Ser 15 20 25 | 474 |
| tgt tgg gca ttc tca gca atc ggt gca gta gaa gga ata aat aag ata Cys Trp Ala Phe Ser Ala Ile Gly Ala Val Glu Gly Ile Asn Lys Ile 30 35 40 | 522 |
| gta aca ggc gaa ctg att tcg tta tca gaa caa gaa ttg gtg gat tgt Val Thr Gly Glu Leu Ile Ser Leu Ser Glu Gln Glu Leu Val Asp Cys 45 50 55 | 570 |
| gat act gga tat aac caa gga tgc aat gga gga ctt atg gac tat gca Asp Thr Gly Tyr Asn Gln Gly Cys Asn Gly Gly Leu Met Asp Tyr Ala 60 65 70 | 618 |
| ttt gag ttc ata atc aac aat ggc ggc att gat tct gat gag gat tac Phe Glu Phe Ile Ile Asn Asn Gly Gly Ile Asp Ser Asp Glu Asp Tyr 75 80 85 90 | 666 |
| cca tac cgt ggt gtt gat ggt aga tgc gac aca tat agg aaa aat gct Pro Tyr Arg Gly Val Asp Gly Arg Cys Asp Thr Tyr Arg Lys Asn Ala 95 100 105 | 714 |
| aaa gtc gtt tct att gat gac tac gaa gat gtt cct gcc tat gat gag Lys Val Val Ser Ile Asp Asp Tyr Glu Asp Val Pro Ala Tyr Asp Glu 110 115 120 | 762 |
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| tct agc tgg gga gag gat ggc tac atc aga tta gaa aga aat ctt gct Ser Ser Trp Gly Glu Asp Gly Tyr Ile Arg Leu Glu Arg Asn Leu Ala 190 195 200 | | | 1002 |
| aac agc aga tca ggc aag tgt gga att gca att gag cca tct tat ccc Asn Ser Arg Ser Gly Lys Cys Gly Ile Ala Ile Glu Pro Ser Tyr Pro 205 210 215 | | | 1050 |
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| tgt ctc agg agc aag aac aac ccc ttt gga gtg aag gca tta agg cgt Cys Leu Arg Ser Lys Asn Asn Pro Phe Gly Val Lys Ala Leu Arg Arg 300 305 310 | | | 1338 |
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Ala Asp Lys Ala Ala Thr Leu Arg Thr Glu Glu Glu Leu Met Ser Met
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Tyr Glu Gln Trp Leu Val Lys His Gly Lys Val Tyr Asn Ala Leu Gly
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Glu Lys Glu Lys Arg Phe Gln Ile Phe Lys Asp Asn Leu Arg Phe Ile
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Asp Asp His Asn Ser Ala Glu Asp Arg Thr Tyr Lys Leu Gly Leu Asn
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Arg Phe Ala Asp Leu Thr Asn Glu Glu Tyr Arg Ala Lys Tyr Leu Gly
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Thr Lys Ile Asp Pro Asn Arg Arg Leu Gly Lys Thr Pro Ser Asn Arg
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Tyr Ala Pro Arg Val Gly Asp Lys Leu Pro Asp Ser Val Asp Trp Arg
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Lys Glu Gly Ala Val Pro Pro Val Lys Asp Gln Gly Gly Cys Gly Ser
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Cys Trp Ala Phe Ser Ala Ile Gly Ala Val Glu Gly Ile Asn Lys Ile
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Val Thr Gly Glu Leu Ile Ser Leu Ser Glu Gln Glu Leu Val Asp Cys
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Asp Thr Gly Tyr Asn Gln Gly Cys Asn Gly Gly Leu Met Asp Tyr Ala
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Phe Glu Phe Ile Ile Asn Asn Gly Gly Ile Asp Ser Asp Glu Asp Tyr
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Pro Tyr Arg Gly Val Asp Gly Arg Cys Asp Thr Tyr Arg Lys Asn Ala
95 100 105

Lys Val Val Ser Ile Asp Asp Tyr Glu Asp Val Pro Ala Tyr Asp Glu
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Leu Ala Leu Lys Lys Ala Val Ala Asn Gln Pro Val Ser Val Ala Ile
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Glu Gly Gly Arg Glu Phe Gln Leu Tyr Val Ser Gly Val Phe Thr
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Gly Arg Cys Gly Thr Ala Leu Asp His Gly Val Val Ala Val Gly Tyr
155 160 165 170

Gly Thr Ala Lys Gly His Asp Tyr Trp Ile Val Arg Asn Ser Trp Gly
175 180 185

Ser Ser Trp Gly Glu Asp Gly Tyr Ile Arg Leu Glu Arg Asn Leu Ala
190 195 200

Asn Ser Arg Ser Gly Lys Cys Gly Ile Ala Ile Glu Pro Ser Tyr Pro
205 210 215

Leu Lys Asn Gly Pro Asn Pro Pro Asn Pro Gly Pro Ser Pro Pro Ser
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Pro Val Lys Pro Pro Asn Val Cys Asp Asn Tyr Tyr Ser Cys Ala Asp
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Ser Ala Thr Cys Cys Cys Ile Phe Glu Phe Gly Asn Ala Cys Phe Glu
255 260 265

Trp Gly Cys Cys Pro Leu Glu Gly Ala Ser Cys Cys Asp Asp His Tyr
270 275 280

Ser Cys Cys Pro Ala Asp Tyr Pro Ile Cys Asn Thr Tyr Ala Gly Thr
285 290 295

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actcattagg caccccaggc tttacacttt atgcttcgg atcgtatgtt gtgtggaatt 120
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tgcctgcagg tcgcccttc gtcttcaaga attccctgt tgacaattaa tcattcaact 180
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